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## THE ENITED STATES PATENT AND TRADEMARK OFFICE

In re application of: Mikal E. Saltveit et al.	)		
Serial No.: <b>09/964,992</b>	Examiner: Not Yet Assigned		
Filed: September 26, 2001	) Art Unit: 1645 )		
For: Characterization of Phenylalanine Ammonia-lyase (PAL) Gene in Wounded Lettuce Tissue	) MARKED-UP VERSION TO SHOW		
	) <u>CHANGES MADE</u>		
	RECEIVED		
	FF		
Assistant Commissioner for Patents Washington, D.C. 20231	TECH CENTER 1600/2900		

Sir:

This marked-up specification (section 'Brief Description of the Drawings', page 5 lines 1-2) accompanies the attached preliminary amendment for the above-identified patent application and is submitted on or before December 26, 2001.

## In the Specification

Figure 9: Sequence comparison of LsPAL1 (SEQ ID NO:1) to sunflower PAL (SEQ ID NO:5) demonstrating the close sequence homology.

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Washington, D.C. 20231 on 12/11/01		

Printed Name: Pube. tra. tison

Mikal E. Saltveit *et al*. Serial No. 09/964,992

Respectfully submitted,

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## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Ammonia-lyase (PAL) Gene in Wounded Lettuce Tissue	) SHOW CHANGES MADE	
For: Characterization of Phenylalanine	SHOW CHANGES MADE	
•	) RED SKETCH OF DRAWING TO	
Filed: September 26, 2001	) Art Unit: 1645 )	
Serial No.: <b>09/964,992</b>	) Am Haire 1645	
	) Examiner: Not Yet Assigned	
In re application of: Mikal E. Saltveit et al.	)	

Assistant Commissioner for Patents Washington, D.C. 20231

Sir

This red sketch of Figure 9 accompanies the attached preliminary amendment for the above-identified patent application and is submitted on or before December 26, 2001.

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Signature: Robert, Ettinic

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+		
1	MENGTHVNG <mark>S</mark> ANG <mark>P</mark> CIKDPLNWGVAAEALTGSHLDEVKKMVGEFRKPVVK MENG <mark>N</mark> HVNGV <mark>V</mark> NE <mark>LCIKDPLNWGVAAEALTGSHLDEVKKMV<mark>A</mark>EFRKPVVK</mark>	Sunflower LsPALI
51	LGGETLTVSQVAGI <mark>S</mark> AAGDG <mark>NM</mark> VKVELSEAARAGVKASSDWVMESMNKGT	Sunflower
51	LGGETLTVSQVAGIAAA <mark>M</mark> D <mark>S</mark> DTVKVELSEAARAGVKASSDWVMESMNKGT	LsPAL1
101	DSYGVTTGFGATSHRRTK <mark>N</mark> GGALQKELIRFLNAGIFGNGTESSHTLPHSA	Sunflower
101	DSYGVTTGFGATSHRRTKQGGALQKELIRFLNAGIFGNGTE <mark>T</mark> SHTLPHSA	LsPAL1
151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLNNNITPCLPLRGTITASGD	Sunflewer
151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLNNNITPCLPLRGTITASGD	LsPAL1
201	LVPLSYIAGLLTGRPNSKAVGPAGEVLNAESAFA <mark>Q</mark> AGVEGGFFELQPKEG	Sunflower
201	LVPLSYIAGLLTGRPNSKAVGP <mark>T</mark> GEVLNAE <mark>K</mark> AFAAAGVEGGFFELQPKEG	LsPAL1
251	LALVNGTAVGSGMASMVLFEANVLALLSEVLSAIFAEVMQGKPEFTDHLT	Sunflower
251	LALVNGTAVGSGMASMVLFDANVLALLSEVLSAIFAEVMQGKPEFTDHLT	LsPAL1
301	HKLKHHPGQIEAAAIMEYILDGSDYVKAAQKVHEMDPLQKPKQDRYALRT	Sunflower
301	HKLKHHPGQIEAAAIMEYILDGSDYVKAAQKVHEMDPLQKPKQDRYALRT	LsPAL1
351	SPQWLGPQIEVIRSATKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	Sunflower
351	SPQWLGPQIEVIRS <mark>S</mark> TKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	LsPAL1
401	GVSMDNTRLAIAAIGKVTIAQFSELVNDFYNNGLPS <mark>H</mark> LSGGRNPSLDSGF	Sunflower
401	GVSMDNTRLAIAAIGK <mark>LMF</mark> AQFSELVNDFYNNGLPSNLSGGRNPSLD <mark>Y</mark> GF	LsPAL1
451	KGGEIAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGLISARKTAEAV	Sunflower
451	KGGEIAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGLISARKTAEAV	LsPAL1
501	DILKLMSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	Sunflower
501	DILKLMSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	LsPAL1
551	LHPSRFCEKDLLRVVDREYVFAYADD <mark>P</mark> CL <mark>T</mark> TYPLMQKLRQVLVDHALNNG	Sunflower
551	LHPSRFCEKDLLRVVDREYVFAY <mark>T</mark> DDVC <mark>S</mark> GTYPLMQKLRQVLVDHALNNG	LsPAL1
601	ETEKNANTSIFQKIATFEDELKA <mark>I</mark> LPKEVE <mark>S</mark> VRVAFENGT <mark>M</mark> SIPNRIKAC	Sunflower
601	ETEKN <mark>I</mark> NTSIFQKIATFE <mark>B</mark> ELK <mark>V</mark> LLPKEVEGVR <mark>IAY</mark> EN <mark>D</mark> TLSIPNRIKAC	LsPAL1
651	RSYPLYRFVREELGG <mark>A</mark>	Sunflewer
651	RSYPLYRFVREELG <mark>R</mark> GFLTGEKVTSPGEEFDRVFTAMCKGQIIDPLLECL	LsPAL1
701	GGWNGEPLPIC	LSPALI